

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/550,797
Source: 1Fw0
Date Processed by STIC: 8/28/06

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/550,797

DATE: 08/28/2006

TIME: 09:26:32

Input Set : F:\seq_210121_609uspc.app.txt
 Output Set: N:\CRF4\08282006\J550797.raw

3 <110> APPLICANT: Zehentner-Wilkinson, Barbara K.
 4 Hayes, Dawn
 5 Houghton, Raymond L.
 7 <120> TITLE OF INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION
 8 AND MONITORING OF LUNG CANCER
 10 <130> FILE REFERENCE: 210121.609USPC
 12 <140> CURRENT APPLICATION NUMBER: US 10/550,797
 C--> 13 <141> CURRENT FILING DATE: 2005-09-22
 15 <160> NUMBER OF SEQ ID NOS: 34
 17 <170> SOFTWARE: Corixa Invention Disclosure Database
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 3951
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapiens
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 27 tggacttc ctgggtgcct taagttcaga actccattc ctgggagctg gaggacatgt 180
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 84 gtttgcattttt ttctactccc atcaaaagcag cttttttttt tattttttttt gttttttttt 3600
 85 atgatagtttta tagccctttaatgccttaa ctaaggaaga aaagatgttta ttctgagttt 3660
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 87 ggagatacta acctttggaa atgattagct ggctctgttttttggtaaaa taagagtctt 3780
 88 taatcctttc tccatcaaga gtacttacc aaggcaggg gaagggggat atagaggc 3840
 89 caaggaaataaaaatcatct ttcatttttta attttactcc ttccttttat tttttttttt 3900
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 93 <211> LENGTH: 943
 94 <212> TYPE: PRT
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 100 Thr Leu Leu Val Ala Leu Ser Ser Glu Leu Pro Phe Leu Gly Ala Gly
 101 20 25 30
 102 Val Gln Leu Gln Asp Asn Gly Tyr Asn Gly Leu Leu Ile Ala Ile Asn
 103 35 40 45

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 105 50 55 60
 106 Ile Thr Glu Ala Ser Phe Tyr Leu Phe Asn Ala Thr Lys Arg Arg Val
 107 65 70 75 80
 108 Phe Phe Arg Asn Ile Lys Ile Leu Ile Pro Ala Thr Trp Lys Ala Asn
 109 85 90 95
 110 Asn Asn Ser Lys Ile Lys Gln Glu Ser Tyr Glu Lys Ala Asn Val Ile
 111 100 105 110
 112 Val Thr Asp Trp Tyr Gly Ala His Gly Asp Asp Pro Tyr Thr Leu Gln
 113 115 120 125
 114 Tyr Arg Gly Cys Gly Lys Glu Gly Lys Tyr Ile His Phe Thr Pro Asn
 115 130 135 140
 116 Phe Leu Leu Asn Asp Asn Leu Thr Ala Gly Tyr Gly Ser Arg Gly Arg
 117 145 150 155 160
 118 Val Phe Val His Glu Trp Ala His Leu Arg Trp Gly Val Phe Asp Glu
 119 165 170 175
 120 Tyr Asn Asn Asp Lys Pro Phe Tyr Ile Asn Gly Gln Asn Gln Ile Lys
 121 180 185 190
 122 Val Thr Arg Cys Ser Ser Asp Ile Thr Gly Ile Phe Val Cys Glu Lys
 123 195 200 205
 124 Gly Pro Cys Pro Gln Glu Asn Cys Ile Ile Ser Lys Leu Phe Lys Glu
 125 210 215 220
 126 Gly Cys Thr Phe Ile Tyr Asn Ser Thr Gln Asn Ala Thr Ala Ser Ile
 127 225 230 235 240
 128 Met Phe Met Gln Ser Leu Ser Ser Val Val Glu Phe Cys Asn Ala Ser
 129 245 250 255
 130 Thr His Asn Gln Glu Ala Pro Asn Leu Gln Asn Gln Met Cys Ser Leu
 131 260 265 270
 132 Arg Ser Ala Trp Asp Val Ile Thr Asp Ser Ala Asp Phe His His Ser
 133 275 280 285
 134 Phe Pro Met Asn Gly Thr Glu Leu Pro Pro Pro Pro Thr Phe Ser Leu
 135 290 295 300
 136 Val Glu Ala Gly Asp Lys Val Val Cys Leu Val Leu Asp Val Ser Ser
 137 305 310 315 320
 138 Lys Met Ala Glu Ala Asp Arg Leu Leu Gln Leu Gln Gln Ala Ala Glu
 139 325 330 335
 140 Phe Tyr Leu Met Gln Ile Val Glu Ile His Thr Phe Val Gly Ile Ala
 141 340 345 350
 142 Ser Phe Asp Ser Lys Gly Glu Ile Arg Ala Gln Leu His Gln Ile Asn
 143 355 360 365
 144 Ser Asn Asp Asp Arg Lys Leu Leu Val Ser Tyr Leu Pro Thr Thr Val
 145 370 375 380
 146 Ser Ala Lys Thr Asp Ile Ser Ile Cys Ser Gly Leu Lys Lys Gly Phe
 147 385 390 395 400
 148 Glu Val Val Glu Lys Leu Asn Gly Lys Ala Tyr Gly Ser Val Met Ile
 149 405 410 415
 150 Leu Val Thr Ser Gly Asp Asp Lys Leu Leu Gly Asn Cys Leu Pro Thr
 151 420 425 430
 152 Val Leu Ser Ser Gly Ser Thr Ile His Ser Ile Ala Leu Gly Ser Ser

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153 435 440 445
 154 Ala Ala Pro Asn Leu Glu Glu Leu Ser Arg Leu Thr Gly Gly Leu Lys
 155 450 455 460
 156 Phe Phe Val Pro Asp Ile Ser Asn Ser Asn Met Ile Asp Ala Phe
 157 465 470 475 480
 158 Ser Arg Ile Ser Ser Gly Thr Gly Asp Ile Phe Gln Gln His Ile Gln
 159 485 490 495
 160 Leu Glu Ser Thr Gly Glu Asn Val Lys Pro His His Gln Leu Lys Asn
 161 500 505 510
 162 Thr Val Thr Val Asp Asn Thr Val Gly Asn Asp Thr Met Phe Leu Val
 163 515 520 525
 164 Thr Trp Gln Ala Ser Gly Pro Pro Glu Ile Ile Leu Phe Asp Pro Asp
 165 530 535 540
 166 Gly Arg Lys Tyr Tyr Thr Asn Asn Phe Ile Thr Asn Leu Thr Phe Arg
 167 545 550 555 560
 168 Thr Ala Ser Leu Trp Ile Pro Gly Thr Ala Lys Pro Gly His Trp Thr
 169 565 570 575
 170 Tyr Thr Leu Asn Asn Thr His His Ser Leu Gln Ala Leu Lys Val Thr
 171 580 585 590
 172 Val Thr Ser Arg Ala Ser Asn Ser Ala Val Pro Pro Ala Thr Val Glu
 173 595 600 605
 174 Ala Phe Val Glu Arg Asp Ser Leu His Phe Pro His Pro Val Met Ile
 175 610 615 620
 176 Tyr Ala Asn Val Lys Gln Gly Phe Tyr Pro Ile Leu Asn Ala Thr Val
 177 625 630 635 640
 178 Thr Ala Thr Val Glu Pro Glu Thr Gly Asp Pro Val Thr Leu Arg Leu
 179 645 650 655
 180 Leu Asp Asp Gly Ala Gly Ala Asp Val Ile Lys Asn Asp Gly Ile Tyr
 181 660 665 670
 182 Ser Arg Tyr Phe Phe Ser Phe Ala Ala Asn Gly Arg Tyr Ser Leu Lys
 183 675 680 685
 184 Val His Val Asn His Ser Pro Ser Ile Ser Thr Pro Ala His Ser Ile
 185 690 695 700
 186 Pro Gly Ser His Ala Met Tyr Val Pro Gly Tyr Thr Ala Asn Gly Asn
 187 705 710 715 720
 188 Ile Gln Met Asn Ala Pro Arg Lys Ser Val Gly Arg Asn Glu Glu
 189 725 730 735
 190 Arg Lys Trp Gly Phe Ser Arg Val Ser Ser Gly Gly Ser Phe Ser Val
 191 740 745 750
 192 Leu Gly Val Pro Ala Gly Pro His Pro Asp Val Phe Pro Pro Cys Lys
 193 755 760 765
 194 Ile Ile Asp Leu Glu Ala Val Lys Val Glu Glu Leu Thr Leu Ser
 195 770 775 780
 196 Trp Thr Ala Pro Gly Glu Asp Phe Asp Gln Gly Gln Ala Thr Ser Tyr
 197 785 790 795 800
 198 Glu Ile Arg Met Ser Lys Ser Leu Gln Asn Ile Gln Asp Asp Phe Asn
 199 805 810 815
 200 Asn Ala Ile Leu Val Asn Thr Ser Lys Arg Asn Pro Gln Gln Ala Gly
 201 820 825 830

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202 Ile Arg Glu Ile Phe Thr Phe Ser Pro Gln Ile Ser Thr Asn Gly Pro
203 835 840 845
204 Glu His Gln Pro Asn Gly Glu Thr His Glu Ser His Arg Ile Tyr Val
205 850 855 860
206 Ala Ile Arg Ala Met Asp Arg Asn Ser Leu Gln Ser Ala Val Ser Asn
207 865 870 875 880
208 Ile Ala Gln Ala Pro Leu Phe Ile Pro Pro Asn Ser Asp Pro Val Pro
209 885 890 895
210 Ala Arg Asp Tyr Leu Ile Leu Lys Gly Val Leu Thr Ala Met Gly Leu
211 900 905 910
212 Ile Gly Ile Ile Cys Leu Ile Ile Val Val Thr His His Thr Leu Ser
213 915 920 925
214 Arg Lys Lys Arg Ala Asp Lys Lys Glu Asn Gly Thr Lys Leu Leu
215 930 935 940

217 <210> SEQ ID NO: 3

218 <211> LENGTH: 785

219 <212> TYPE: DNA

220 <213> ORGANISM: Homo sapiens

222 <400> SEQUENCE: 3

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225 ttttcttg ccacggccgc agccgcggcg gcccgaccc cccgaccc 180
226 gccgacgcgc agcagcagca gcagcagcag caggcgccgc agctgagacc ggccgcgc 240
227 ggccagccct cagggggcgcc tcacaagtca ggcggccaa aagtcaagcg acagcgctcg 300
228 tcttcggcccg aactgatgcg ctgcaaaacgc cggctcaact tcagcggctt tggctacagc 360
229 ctggccgcgc agcagccggc cgccgtggcg cgccgcaacg agcgcgagcg caaccgcgtc 420
230 aagtgggtca acctgggtt tgccaccctt cgggagcagc tccccaaacgg cgccggccaa 480
231 aagaagatga gtaagggtgga gacactgcgc tcggcggtcg agtacatccg cgcgtgcag 540
232 cagctgctgg acgagcatga cgcgggtgagc gcccgttcc aggcaggcgt cctgtcgccc 600
233 accatctccc ccaactactc caacgacttg aactccatgg ccggctcgcc ggtctcatcc 660
234 tactcgtcggt acgagggtctc ttacgaccgg ctcagccccg aggagcagga gttctcgac 720
235 ttcaccaact ggttctgagg ggctcggcct ggtcaggccc tggtgcaat ggactttgga 780
236 agcag 785

238 <210> SEQ ID NO: 4

239 <211> LENGTH: 236

240 <212> TYPE: PRT

241 <213> ORGANISM: Homo sapiens

243 <400> SEQUENCE: 4

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248 Ala Thr Ala Gln
249 35 40 45
250 Ser Ala Gln Ala Pro
251 50 55 60
252 Gln Leu Arg Pro Ala Ala Asp Gly Gln Pro Ser Gly Gly His Lys
253 65 70 75 80
254 Ser Ala Pro Lys Gln Val Lys Arg Gln Arg Ser Ser Ser Pro Glu Leu

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date